

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/529,613
Source: JFWP
Date Processed by STIC: 07/05/2006

ENTERED



IFWP

RAW SEQUENCE LISTING

DATE: 07/05/2006

PATENT APPLICATION: US/10/529,613

TIME: 14:13:48

Input Set : A:\SEQLIST.TXT

Output Set: N:\CRF4\07052006\J529613.raw

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4 <110> APPLICANT: Kudaravalli, Sridar
5      Torres, Rosarelis
6      Wolfgang, Curt
7      Polymeropoulos, Mihael
9 <120> TITLE OF INVENTION: METHODS TO PREDICT CHOLESTEROL
10     ELEVATIONS DURING IMMUNOSUPPRESSANT THERAPY
13 <130> FILE REFERENCE: DC/4-32702A
15 <140> CURRENT APPLICATION NUMBER: 10/529,613
C--> 16 <141> CURRENT FILING DATE: 2005-03-30
18 <150> PRIOR APPLICATION NUMBER: PCT/EP03/10798
19 <151> PRIOR FILING DATE: 2003-09-29
21 <150> PRIOR APPLICATION NUMBER: 60/415, 123
22 <151> PRIOR FILING DATE: 2002-09-30
24 <160> NUMBER OF SEQ ID NOS: 8
26 <170> SOFTWARE: FastSEQ for Windows Version 4.0
28 <210> SEQ ID NO: 1
29 <211> LENGTH: 20
30 <212> TYPE: DNA
31 <213> ORGANISM: Artificial Sequence
33 <220> FEATURE:
34 <221> NAME/KEY: primer_bind
35 <222> LOCATION: (1)...(20)
36 <223> OTHER INFORMATION: IL-1 (-511)-forward primer
38 <223> OTHER INFORMATION: Synthetic oligonucleotide
W--> 40 <400> 1
41 gcagagctca tctggcattg
43 <210> SEQ ID NO: 2
44 <211> LENGTH: 20
45 <212> TYPE: DNA
46 <213> ORGANISM: Artificial Sequence
48 <220> FEATURE:
49 <221> NAME/KEY: primer_bind
50 <222> LOCATION: (1)...(20)
51 <223> OTHER INFORMATION: IL-1 (-511)-reverse primer
53 <223> OTHER INFORMATION: Synthetic oligonucleotide
W--> 55 <400> 2
56 tatgtgggac aaagtggaag
58 <210> SEQ ID NO: 3
59 <211> LENGTH: 22
60 <212> TYPE: DNA
C--> 61 <213> ORGANISM: Artifical Sequence
63 <220> FEATURE:
64 <221> NAME/KEY: primer_bind

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65 <222> LOCATION: (1)...(22)
66 <223> OTHER INFORMATION: IL-1 (-31)-forward primer
68 <223> OTHER INFORMATION: Synthetic oligonucleotide
W--> 70 <400> 3
71 gcacaacgat tgtcaggaaa ac 22
73 <210> SEQ ID NO: 4
74 <211> LENGTH: 22
75 <212> TYPE: DNA
76 <213> ORGANISM: Artificial Sequence
78 <220> FEATURE:
79 <221> NAME/KEY: primer_bind
80 <222> LOCATION: (1)...(22)
81 <223> OTHER INFORMATION: IL-1 (-31)-reverse primer
83 <223> OTHER INFORMATION: Synthetic oligonucleotide
W--> 85 <400> 4
86 atgcatacac acaaagaggc ag 22
88 <210> SEQ ID NO: 5
89 <211> LENGTH: 55
90 <212> TYPE: DNA
91 <213> ORGANISM: Homo sapiens
93 <220> FEATURE:
94 <221> NAME/KEY: allele
95 <222> LOCATION: (1)...(55)
96 <223> OTHER INFORMATION: Nucleotide sequence surrounding the (-511) IL-1
97 polymorphism, allele 1
99 <400> SEQUENCE: 5
100 ctgcaattga cagagagctc ccgaggcaga gaacagcacc caaggtagag accca 55
102 <210> SEQ ID NO: 6
103 <211> LENGTH: 55
104 <212> TYPE: DNA
105 <213> ORGANISM: Homo sapiens
107 <220> FEATURE:
108 <221> NAME/KEY: allele
109 <222> LOCATION: (1)...(55)
110 <223> OTHER INFORMATION: Nucleotide sequence surrounding the (-511) IL-1
111 polymorphism, allele 2
113 <400> SEQUENCE: 6
114 ctgcaattga cagagagctc ctgaggcaga gaacagcacc caaggtagag accca 55
116 <210> SEQ ID NO: 7
117 <211> LENGTH: 63
118 <212> TYPE: DNA
119 <213> ORGANISM: Homo sapiens
121 <220> FEATURE:
122 <221> NAME/KEY: allele
123 <222> LOCATION: (1)...(63)
124 <223> OTHER INFORMATION: Nucleotide sequence surrounding the (-31) IL-1
125 polymorphism, allele 1
127 <400> SEQUENCE: 7
128 tcctacttct gcttttgaaa gccataaaaa cagcgaggga gaaactggca gataccaaac 60

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129 ctc 63
131 <210> SEQ ID NO: 8
132 <211> LENGTH: 63
133 <212> TYPE: DNA
134 <213> ORGANISM: Homo sapiens
136 <220> FEATURE:
137 <221> NAME/KEY: allele
138 <222> LOCATION: (1)...(63)
139 <223> OTHER INFORMATION: Nucleotide sequence surrounding the (-31) IL-1
140 polymorphism, allele 2
142 <400> SEQUENCE: 8
143 tcctacttct gcttttgaaa gctataaaaa cagcgaggga gaaactggca gataccaaac 60
144 ctc 63
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VERIFICATION SUMMARY

DATE: 07/05/2006

PATENT APPLICATION: US/10/529,613

TIME: 14:13:50

Input Set : A:\SEQLIST.TXT

Output Set: N:\CRF4\07052006\J529613.raw

L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:40 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:55 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2
L:61 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3
L:70 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:85 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4